

Atty. Docket No. 8062-1023

PATENTS

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re application of

Fumihide NISHIO

Confirmation No. 9947

Serial No. 10/501,671
(PCT/JP03/00339)

PCT/DO/EO

Filed July 16, 2004

HIGH-CONCENTRATION PREPARATION
OF SOLUBLE THROMBOMODULIN

RESPONSE TO NOTIFICATION OF MISSING REQUIREMENTS

Commissioner for Patents
P.O. Box 1450
Alexandria, VA 22313-1450

June 28, 2005

Sir:

In response to the accompanying Notification of Missing Requirements under 35 U.S.C. § 371 mailed March 31, 2005, we enclose herewith the executed Declaration, which was omitted at the time of filing the application. The required 37 C.F.R. § 1.492(e) surcharge was paid at the time of filing the application. Also enclosed is a Supplemental Application Data Sheet.

Attached herewith is a Statement to Support Filing and

Docket No. 8062-1023
Appln. No. 10/501,671

Submission in Accordance with 37 C.F.R. §§ 1.821-1.825 with the
Sequence Listing in paper and disk formats.

Respectfully submitted,

YOUNG & THOMPSON

A handwritten signature in black ink, appearing to read "Michael Piziali", is written over a horizontal line.

Michael Piziali, Reg. No. 46,997
745 South 23rd Street
Arlington, VA 22202
Telephone (703) 521-2297

MP:fb



UNITED STATES PATENT AND TRADEMARK OFFICE

UNITED STATES DEPARTMENT OF COMMERCE
 United States Patent and Trademark Office
 Address: COMMISSIONER FOR PATENTS
 P.O. Box 1450
 Alexandria, Virginia 22313-1450
 www.uspto.gov

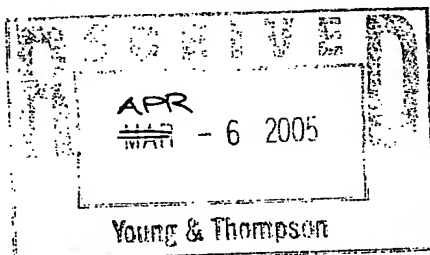
| | | |
|-----------------------------|-----------------------|------------------|
| U.S. APPLICATION NUMBER NO. | FIRST NAMED APPLICANT | ATTY. DOCKET NO. |
| 10/501,671 | Fumihide Nishio | 8062-1023 |

| |
|-------------------------------|
| INTERNATIONAL APPLICATION NO. |
|-------------------------------|

PCT/JP03/00339

| | |
|------------------|---------------|
| I.A. FILING DATE | PRIORITY DATE |
| 01/17/2003 | 01/18/2002 |

00466
 YOUNG & THOMPSON
 745 SOUTH 23RD STREET
 2ND FLOOR
 ARLINGTON, VA 22202



CONFIRMATION NO. 9947

371 FORMALITIES LETTER



OC000000015512526

Date Mailed: 03/31/2005

NOTIFICATION OF MISSING REQUIREMENTS UNDER 35 U.S.C. 371 IN THE UNITED STATES DESIGNATED/ELECTED OFFICE (DO/EO/US)

The following items have been submitted by the applicant or the IB to the United States Patent and Trademark Office as a Designated / Elected Office (37 CFR 1.495).

- Copy of the International Application filed on 07/16/2004
- Copy of the International Search Report filed on 07/16/2004
- Copy of IPE Report filed on 07/16/2004
- Preliminary Amendments filed on 07/16/2004
- Information Disclosure Statements filed on 07/16/2004
- Biochemical Sequence Diskette filed on 07/16/2004
- Biochemical Sequence Listing filed on 07/16/2004
- Request for Immediate Examination filed on 07/16/2004
- Copy of references cited in ISR filed on 07/16/2004
- U.S. Basic National Fees filed on 07/16/2004
- Priority Documents filed on 07/16/2004

The following items **MUST** be furnished within the period set forth below in order to complete the requirements for acceptance under 35 U.S.C. 371:

- Oath or declaration of the inventors, in compliance with 37 CFR 1.497(a) and (b), identifying the application by the International application number and international filing date.
- This application clearly fails to comply with the requirements of 37 CFR. 1.821-1.825. Applicant's attention is directed to the final rulemaking notice published at 55 FR 18230 (May 1, 1990), and 1114 OG 29 (May 15, 1990). If the effective filing date is on or after July 1, 1998, see the final rulemaking notice published at 63 FR 29620 (June 1, 1998) and 1211 OG 82 (June 23, 1998). If the effective filing date is on or after September 8, 2000, see the final rulemaking notice published in the Federal Register at 65 FR 54604 (September 8, 2000) and 1238 OG 145 (September 19, 2000). Applicant must provide an initial computer readable form (CRF) copy of the "Sequence Listing", an initial paper or compact disc copy of the "Sequence Listing", as well as an amendment directing its entry into the application. Applicant must also

provide a statement that the content of the sequence listing information recorded in computer readable form is identical to the written (on paper or compact disc) sequence listing and, where applicable, includes no new matter, as required by 37 CFR 1.821(e), 1.821(f), 1.821(g), 1.825(b), or 1.825(d). If applicant desires the sequence listing in the instant application to be identical with that of another application on file in the U.S. Patent and Trademark Office, such request in accordance with 37 CFR 1.821(e) may be submitted in lieu of a new CRF.

- This application does not contain a statement that the content of the sequence listing information recorded in computer readable form is identical to the written (on paper or compact disc) sequence listing and, where applicable, includes no new matter, as required by 37 CFR 1.821(e), 1.821(f), 1.821(g), 1.825(b), or 1.825(d). Applicant must provide such statement. If the effective filing date is on or after September 8, 2000, see the final rulemaking notice published in the Federal Register at 65 FR 54604 (September 8, 2000) and 1238 OG 145 (September 19, 2000).

ALL OF THE ITEMS SET FORTH ABOVE MUST BE SUBMITTED WITHIN TWO (2) MONTHS FROM THE DATE OF THIS NOTICE OR BY 32 MONTHS FROM THE PRIORITY DATE FOR THE APPLICATION, WHICHEVER IS LATER. FAILURE TO PROPERLY RESPOND WILL RESULT IN ABANDONMENT.

The time period set above may be extended by filing a petition and fee for extension of time under the provisions of 37 CFR 1.136(a).

For questions regarding compliance to 37 CFR 1.821-1.825 requirements, please contact:

- For Rules Interpretation, call (571) 272-0951
- For Patentin Software Program Help, call Patent EBC at 1-866-217-9197 or directly at 703-305-3028 / 703-308-6845 between the hours of 6 a.m. and 12 midnight, Monday through Friday, EST.
- Send e-mail correspondence for Patentin Software Program Help @ ebc@uspto.gov

Applicant is reminded that any communications to the United States Patent and Trademark Office must be mailed to the address given in the heading and include the U.S. application no. shown above (37 CFR 1.5)

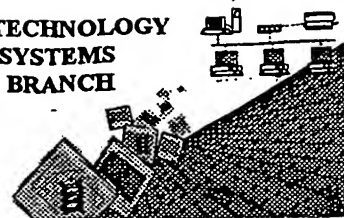
*A copy of this notice **MUST** be returned with the response.*

CHRISTINE S WASHINGTON

Telephone: (703) 308-9140 EXT 228

PART 1 - ATTORNEY/APPLICANT COPY

| U.S. APPLICATION NUMBER NO. | INTERNATIONAL APPLICATION NO. | ATTY. DOCKET NO. |
|-----------------------------|-------------------------------|------------------|
| 10/501,671 | PCT/JP03/00339 | 8062-1023 |



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/501,671
Source: PG/10
Date Processed by STIC: 7/23/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 4.2 PROGRAM. ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebs/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop-Sequence, Crystal Plaza Two, Lo
Room 1B03, Arlington, VA 22202

Revised 05/17/04

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10/501,671

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 ☐ Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 ☐ Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 ☒ Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 ☐ Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 ☐ Variable Length Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 ☐ PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 ☐ Skipped Sequences
 (OLD RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 ☐ Skipped Sequences
 (NEW RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 ☐ Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 ☐ Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 ☐ Use of <220> Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 ☐ PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 ☐ Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



PCT

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/501,671

DATE: 07/23/2004
TIME: 15:23:45

Input Set : A:\PTO.FG.txt
Output Set: N:\CRF4\07232004\J501671.raw

3 <110> APPLICANT: Asahi Kasei Kabushiki Kaisha
W--> 4 <120> TITLE OF INVENTION: High-concentration preparation of soluble thrombomodulin
6 <130> FILE REFERENCE: ASAHI-33
C--> 8 <140> CURRENT APPLICATION NUMBER: US/10/501,671
C--> 8 <141> CURRENT FILING DATE: 2004-07-16
8 <150> PRIOR APPLICATION NUMBER: JP2002-009951
9 <151> PRIOR FILING DATE: 2002-01-18
11 <160> NUMBER OF SEQ ID NOS: 9

ERRORED SEQUENCES

13 <210> SEQ ID NO: 1
14 <211> LENGTH: 516
15 <212> TYPE: PRT
16 <213> ORGANISM: Artificial sequence
18 <220> FEATURE:
19 <223> OTHER INFORMATION: Partial amino acid sequence of human-originated soluble
20 thrombomodulin
22 <400> SEQUENCE: 1

23 Met Leu Gly Val Leu Val Leu Gly Ala Leu Ala Leu Ala Gly Leu Gly
E--> 24 1 5 5 10 10 15
25 Phe Pro Ala Pro Ala Glu Pro Gln Pro Gly Gly Ser Gln Cys Val Glu
E--> 26 20 20 25 25 30
27 His Asp Cys Phe Ala Leu Tyr Pro Gly Pro Ala Thr Phe Leu Asn Ala
E--> 28 35 40 45
29 Ser Gln Ile Cys Asp Gly Leu Arg Gly His Leu Met Thr Val Arg Ser
E--> 30 50 55 60
31 Ser Val Ala Ala Asp Val Ile Ser Leu Leu Asn Gly Asp Gly Gly
E--> 32 65 70 75
33 Val Gly Arg Arg Arg Leu Trp Ile Gly Leu Gln Leu Pro Pro Gly Cys
E--> 34 85 90 95
35 Gly Asp Pro Lys Arg Leu Gly Pro Leu Arg Gly Phe Gln Trp Val Thr
E--> 36 100 105 110
37 Gly Asp Asn Asn Thr Ser Tyr Ser Arg Trp Ala Arg Leu Asp Leu Asn
E--> 38 115 120 125
39 Gly Ala Pro Leu Cys Gly Pro Leu Cys Val Ala Val Ser Ala Ala Glu
E--> 40 130 135 140
41 Ala Thr Val Pro Ser Glu Pro Ile Trp Glu Glu Gln Gln Cys Glu Val
E--> 42 145 150 155 160
43 Lys Ala Asp Gly Phe Leu Cys Glu Phe His Phe Pro Ala Thr Cys Arg
E--> 44 165 170 175
45 Pro Leu Ala Val Glu Pro Gly Ala Ala Ala Ala Val Ser Ile Thr

misaligned amino acid numbers
15 (see item 3 on Error summary sheet)
80

see p. 2

DATE: 07/23/2004

TIME: 15:23:45

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\07232004\J501671.raw

| | | | | |
|---|-----|-----|-----|-----|
| E--> 46 | 180 | 185 | 190 | |
| 47 Tyr Gly Thr Pro Phe Ala Ala Arg Gly Ala Asp Phe Gln Ala Leu Pro | | | | |
| E--> 48 | 195 | 200 | 205 | |
| 49 Val Gly Ser Ser Ala Ala Val Ala Pro Leu Gly Leu Gln Leu Met Cys | | | | |
| E--> 50 | 210 | 215 | 220 | |
| 51 Thr Ala Pro Pro Gly Ala Val Gln Gly His Trp Ala Arg Glu Ala Pro | | | | |
| E--> 52 | 225 | 230 | 235 | 240 |
| 53 Gly Ala Trp Asp Cys Ser Val Glu Asn Gly Gly Cys Glu His Ala Cys | | | | |
| E--> 54 | 245 | 250 | 255 | |
| 55 Asn Ala Ile Pro Gly Ala Pro Arg Cys Gln Cys Pro Ala Gly Ala Ala | | | | |
| E--> 56 | 260 | 265 | 270 | |
| 57 Leu Gln Ala Asp Gly Arg Ser Cys Thr Ala Ser Ala Thr Gln Ser Cys | | | | |
| E--> 58 | 275 | 280 | 285 | |
| 59 Asn Asp Leu Cys Glu His Phe Cys Val Pro Asn Pro Asp Gln Pro Gly | | | | |
| E--> 60 | 290 | 295 | 300 | |
| 61 Ser Tyr Ser Cys Met Cys Glu Thr Gly Tyr Arg Leu Ala Ala Asp Gln | | | | |
| E--> 62 | 305 | 310 | 315 | 320 |
| 63 His Arg Cys Glu Asp Val Asp Asp Cys Ile Leu Glu Pro Ser Pro Cys | | | | |
| E--> 64 | 325 | 330 | 335 | |
| 65 Pro Gln Arg Cys Val Asn Thr Gln Gly Gly Phe Glu Cys His Cys Tyr | | | | |
| E--> 66 | 340 | 345 | 350 | |
| 67 Pro Asn Tyr Asp Leu Val Asp Gly Glu Cys Val Glu Pro Val Asp Pro | | | | |
| E--> 68 | 355 | 360 | 365 | |
| 69 Cys Phe Arg Ala Asn Cys Glu Tyr Gln Cys Gln Pro Leu Asn Gln Thr | | | | |
| E--> 70 | 370 | 375 | 380 | |
| 71 Ser Tyr Leu Cys Val Cys Ala Glu Gly Phe Ala Pro Ile Pro His Glu | | | | |
| E--> 72 | 385 | 390 | 395 | 400 |
| 73 Pro His Arg Cys Gln Met Phe Cys Asn Gln Thr Ala Cys Pro Ala Asp | | | | |
| E--> 74 | 405 | 410 | 415 | |
| 75 Cys Asp Pro Asn Thr Gln Ala Ser Cys Glu Cys Pro Glu Gly Tyr Ile | | | | |
| E--> 76 | 420 | 425 | 430 | |
| 77 Leu Asp Asp Gly Phe Ile Cys Thr Asp Ile Asp Glu Cys Glu Asn Gly | | | | |
| E--> 78 | 435 | 440 | 445 | |
| 79 Gly Phe Cys Ser Gly Val Cys His Asn Leu Pro Gly Thr Phe Glu Cys | | | | |
| E--> 80 | 450 | 455 | 460 | |
| 81 Ile Cys Gly Pro Asp Ser Ala Leu Val Arg His Ile Gly Thr Asp Cys | | | | |
| E--> 82 | 465 | 470 | 475 | 480 |
| 83 Asp Ser Gly Lys Val Asp Gly Gly Asp Ser Gly Ser Gly Glu Pro Pro | | | | |
| E--> 84 | 485 | 490 | 495 | |
| 85 Pro Ser Pro Thr Pro Gly Ser Thr Leu Thr Pro Pro Ala Val Gly Leu | | | | |
| E--> 86 | 500 | 505 | 510 | |
| 87 Val His Ser Gly | | | | |
| E--> 88 | 515 | | | |
| 90 <210> SEQ ID NO: 2 | | | | |
| 91 <211> LENGTH: 1548 | | | | |
| 92 <212> TYPE: DNA | | | | |
| 93 <213> ORGANISM: Artificial sequence | | | | |
| 95 <220> FEATURE: | | | | |
| 96 <223> OTHER INFORMATION: Partial base sequence of human-originated soluble | | | | |

same
error.

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/501,671

DATE: 07/23/2004

TIME: 15:23:45

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\07232004\J501671.raw

```

97      thrombomodulin gene
99 <400> SEQUENCE: 2
100 atgcttgggg tcttggtcct tggcgcgctg gccctggcgg gccctgggggtt ccccgccaccc 60
101 gcagagccgc agccgggtgg cagccagtgc gtcgagcagc actgcttcgc gctctacccg 120
102 ggccccgcga ccttcctcaa tgccagtcat atctgcgacg gactgcgggg ccacctaata 180
103 acagtgcgct cctcgggtggc tgccgatgtc atttccttgc tactgaacgg cgacggcggc 240
104 gttggccgcc ggcgccctctg gatcggcctg cagctgccac ccggctgcgg cgaccccaag 300
105 cgctcggggc ccctgcgcgg cttccagtgg gttacggggag acaacaacac cagctatagc 360
106 aggtgggcac ggctcgacct caatggggct cccctctgcg gcccggttggtg cgtcgtgtgc 420
107 tccgctgctg agggcactgt gcccagcgag ccgatctggg aggagcagca gtgcgaagtg 480
108 aaggccgatg gcttcctctg cgagttccac ttcccagcca cctgcaggcc actggctgtg 540
109 gagcccgccg ccgcggtctg cgccgtctcg atcacctacg gcaccccggt cgcgcccgcc 600
110 ggagcgggact tccaggcgct gccgggtgggc agctccgcgg cgggtggctcc cctcggctta 660
111 cagctaatagt gcaccgcgcc gccgggagcg gtccaggggc actggggccag ggaggcgccg 720
112 ggcgcttggg actgcagcgt ggagaacggc ggctgcgagc acgcgtgcaa tgcgatccct 780
113 ggggctcccc gctgccagtg cccagccggc gccgccttgc aggcagacgg gcgctcctgc 840
114 accgcatccg cgacgcagtc ctgcaacgac ctctgcgagc atttctgcgt tcccaacccc 900
115 gaccagccgg gctcctactc gtgcatgtgc gagaccggct accggctggc ggccgaccaa 960
116 caccggtgcg aggacgtgga tgactgcata ctggagccca gtccgtgtcc gcagcgtgtg 1020
117 gtcaacacac aggtgggctt cgagtgccac tgctacccta actacgacct ggtggacggc 1080
118 gagtgtgtgg agcccgtgga cccgtgcttc agagccaact gcgagtacca gtgccagccc 1140
119 ctgaacaaaa ctagtacct ctgcgtctgc gccgagggtt tgcgcccatt tcccacagag 1200
120 ccgcacaggt gccagatgtt ttgcaaccag actgcctgtc cagccgactg cgaacccaac 1260
121 acccaggcta gctgtgagtg ccctgaaggc tacatcctgg acgacgggtt catctgcacg 1320
122 gacatcgacg agtgcgaaaa cggcggttct tgctccgggg tgtgccacaa cctccccggg 1380
E--> 123 accttcgagt gcatctgcgg gcccgactcg gcccttgctc gccacattgg caccgac 1440/1439
E--> 124 gactccggca aggtggacgg tggcgacagc ggctctggcg agcccccgcc cagcccgacg 1500/1499
E--> 125 cccggtcca ccttgactcc tccggccgtg gggctcgtgc attcgggc 1547 1548
127 <210> SEQ ID NO: 3
128 <211> LENGTH: 132
129 <212> TYPE: PRT
130 <213> ORGANISM: Artificial sequence
132 <220> FEATURE:
133 <223> OTHER INFORMATION: Partial amino acid sequence of human-originated soluble
134      thrombomodulin
E--> 136 <400> SEQUENCE: 8 3 ← change to
137 Met Leu Gly Val Leu Val Leu Gly Ala Leu Ala Leu Ala Gly Leu Gly
E--> 138 1 5 10 15
139 Phe Pro Asp Pro Cys Phe Arg Ala Asn Cys Glu Tyr Gln Cys Gln Pro
E--> 140 20 25 30
141 Leu Asn Gln Thr Ser Tyr Leu Cys Val Cys Ala Glu Gly Phe Ala Pro
E--> 142 35 40 45
143 Ile Pro His Glu Pro His Arg Cys Gln Met Phe Cys Asn Gln Thr Ala
E--> 144 50 55 60
145 Cys Pro Ala Asp Cys Asp Pro Asn Thr Gln Ala Ser Cys Glu Cys Pro
E--> 146 65 70 75 80
147 Glu Gly Tyr Ile Leu Asp Asp Gly Phe Ile Cys Thr Asp Ile Asp Glu
E--> 148 85 90 95
149 Cys Glu Asn Gly Gly Phe Cys Ser Gly Val Cys His Asn Leu Pro Gly

```

This is a group of 9 positions, not 10.

FYI: periods are invalid in nucleic acid. disregard

misaligned amino acid numbers
see p. 4

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/501,671

DATE: 07/23/2004

TIME: 15:23:45

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\07232004\J501671.raw

misaligned
numbers

100 105 110

E--> 150
151 Thr Phe Glu Cys Ile Cys Gly Pro Asp Ser Ala Leu Val Arg His Ile
E--> 152 115 120 125
153 Gly Thr Asp Cys
E--> 154 130
174 <210> SEQ ID NO: 5
175 <211> LENGTH: 516
176 <212> TYPE: PRT
177 <213> ORGANISM: Artificial sequence
179 <220> FEATURE:
180 <223> OTHER INFORMATION: Partial amino acid sequence of human-originated soluble
181 thrombomodulin
183 <400> SEQUENCE: 5
184 Met Leu Gly Val Leu Val Leu Gly Ala Leu Ala Leu Ala Gly Leu Gly
E--> 185 1 5 10 15
186 Phe Pro Ala Pro Ala Glu Pro Gln Pro Gly Gly Ser Gln Cys Val Glu
E--> 187 20 25 30
188 His Asp Cys Phe Ala Leu Tyr Pro Gly Pro Ala Thr Phe Leu Asn Ala
E--> 189 35 40 45
190 Ser Gln Ile Cys Asp Gly Leu Arg Gly His Leu Met Thr Val Arg Ser
E--> 191 50 55 60
192 Ser Val Ala Ala Asp Val Ile Ser Leu Leu Leu Asn Gly Asp Gly Gly
E--> 193 65 70 75 80
194 Val Gly Arg Arg Arg Leu Trp Ile Gly Leu Gln Leu Pro Gly Cys
E--> 195 85 90 95
196 Gly Asp Pro Lys Arg Leu Gly Pro Leu Arg Gly Phe Gln Trp Val Thr
E--> 197 100 105 110
198 Gly Asp Asn Asn Thr Ser Tyr Ser Arg Trp Ala Arg Leu Asp Leu Asn
E--> 199 115 120 125
200 Gly Ala Pro Leu Cys Gly Pro Leu Cys Val Ala Val Ser Ala Ala Glu
E--> 201 130 135 140
202 Ala Thr Val Pro Ser Glu Pro Ile Trp Glu Glu Gln Gln Cys Glu Val
E--> 203 145 150 155 160
204 Lys Ala Asp Gly Phe Leu Cys Glu Phe His Phe Pro Ala Thr Cys Arg
E--> 205 165 170 175
206 Pro Leu Ala Val Glu Pro Gly Ala Ala Ala Ala Val Ser Ile Thr
E--> 207 180 185 190
208 Tyr Gly Thr Pro Phe Ala Ala Arg Gly Ala Asp Phe Gln Ala Leu Pro
E--> 209 195 200 205
210 Val Gly Ser Ser Ala Ala Val Ala Pro Leu Gly Leu Gln Leu Met Cys
E--> 211 210 215 220
212 Thr Ala Pro Pro Gly Ala Val Gln Gly His Trp Ala Arg Glu Ala Pro
E--> 213 225 230 235 240
214 Gly Ala Trp Asp Cys Ser Val Glu Asn Gly Gly Cys Glu His Ala Cys
E--> 215 245 250 255
216 Asn Ala Ile Pro Gly Ala Pro Arg Cys Gln Cys Pro Ala Gly Ala Ala
E--> 217 260 265 270
218 Leu Gln Ala Asp Gly Arg Ser Cys Thr Ala Ser Ala Thr Gln Ser Cys
E--> 219 275 280 285

misaligned
amino acid
numbers

P.5

RAW SEQUENCE LISTING

DATE: 07/23/2004

PATENT APPLICATION: US/10/501,671

TIME: 15:23:45

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\07232004\J501671.raw

same error

```

220 Asn Asp Leu Cys Glu His Phe Cys Val Pro Asn Pro Asp Gln Pro Gly
E--> 221      290                      295                      300
222 Ser Tyr Ser Cys Met Cys Glu Thr Gly Tyr Arg Leu Ala Ala Asp Gln
E--> 223 305                      310                      315                      320
224 His Arg Cys Glu Asp Val Asp Asp Cys Ile Leu Glu Pro Ser Pro Cys
E--> 225                      325                      330                      335
226 Pro Gln Arg Cys Val Asn Thr Gln Gly Gly Phe Glu Cys His Cys Tyr
E--> 227                      340                      345                      350
228 Pro Asn Tyr Asp Leu Val Asp Gly Glu Cys Val Glu Pro Val Asp Pro
E--> 229                      355                      360                      365
230 Cys Phe Arg Ala Asn Cys Glu Tyr Gln Cys Gln Pro Leu Asn Gln Thr
E--> 231      370                      375                      380
232 Ser Tyr Leu Cys Val Cys Ala Glu Gly Phe Ala Pro Ile Pro His Glu
E--> 233 385                      390                      395                      400
234 Pro His Arg Cys Gln Met Phe Cys Asn Gln Thr Ala Cys Pro Ala Asp
E--> 235                      405                      410                      415
236 Cys Asp Pro Asn Thr Gln Ala Ser Cys Glu Cys Pro Glu Gly Tyr Ile
E--> 237                      420                      425                      430
238 Leu Asp Asp Gly Phe Ile Cys Thr Asp Ile Asp Glu Cys Glu Asn Gly
E--> 239      435                      440                      445
240 Gly Phe Cys Ser Gly Val Cys His Asn Leu Pro Gly Thr Phe Glu Cys
E--> 241      450                      455                      460
242 Ile Cys Gly Pro Asp Ser Ala Leu Ala Arg His Ile Gly Thr Asp Cys
E--> 243 465                      470                      475                      480
244 Asp Ser Gly Lys Val Asp Gly Gly Asp Ser Gly Ser Gly Glu Pro Pro
E--> 245                      485                      490                      495
246 Pro Ser Pro Thr Pro Gly Ser Thr Leu Thr Pro Pro Ala Val Gly Leu
E--> 247                      500                      505                      510
248 Val His Ser Gly
E--> 249      515
288 <210> SEQ ID NO: 7
289 <211> LENGTH: 132
290 <212> TYPE: PRT
291 <213> ORGANISM: Artificial sequence
W--> 292 <220> FEATURE:
293 <223> OTHER INFORMATION: Partial amino acid sequence of human-originated soluble
294      thrombomodulin
296 <400> SEQUENCE: 7
297 Met Leu Gly Val Leu Val Leu Gly Ala Leu Ala Leu Ala Gly Leu Gly
E--> 298      1          5          10          15
299 Phe Pro Asp Pro Cys Phe Arg Ala Asn Cys Glu Tyr Gln Cys Gln Pro
E--> 300          20          25          30
301 Leu Asn Gln Thr Ser Tyr Leu Cys Val Cys Ala Glu Gly Phe Ala Pro
E--> 302          35          40          45
303 Ile Pro His Glu Pro His Arg Cys Gln Met Phe Cys Asn Gln Thr Ala
E--> 304          50          55          60
305 Cys Pro Ala Asp Cys Asp Pro Asn Thr Gln Ala Ser Cys Glu Cys Pro
E--> 306          65          70          75          80
307 Glu Gly Tyr Ile Leu Asp Asp Gly Phe Ile Cys Thr Asp Ile Asp Glu

```

misaligned amino acid numbers
P. 6

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/501,671

DATE: 07/23/2004

TIME: 15:23:45

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\07232004\J501671.raw

mesalised amino acid

95 numbers

```

E--> 308                                     85                                     90
      309 Cys Glu Asn Gly Gly Phe Cys Ser Gly Val Cys His Asn Leu Pro Gly
E--> 310                                     100                                     105                                     110
      311 Thr Phe Glu Cys Ile Cys Gly Pro Asp Ser Ala Leu Ala Arg His Ile
E--> 312                                     115                                     120                                     125
      313 Gly Thr Asp Cys
E--> 314                                     130
      334 <210> SEQ ID NO: 9
      335 <211> LENGTH: 21
      336 <212> TYPE: DNA
      337 <213> ORGANISM: Artificial sequence
      339 <220> FEATURE:
      340 <223> OTHER INFORMATION: Synthetic DNA for mutation
      342 <400> SEQUENCE: 9
      343 aatgtggcgg gcaagggccg a
E--> 349 1/12

```

delete

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/501,671

DATE: 07/23/2004
TIME: 15:23:46

Input Set : A:\PTO.FG.txt
Output Set: N:\CRF4\07232004\J501671.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 32
Seq#:3; Line(s) 146
Seq#:4; Line(s) 172
Seq#:7; Line(s) 306
Seq#:8; Line(s) 332

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/501,671

DATE: 07/23/2004

TIME: 15:23:46

Input Set : A:\PTO.FG.txt

Output Set: N:\CRF4\07232004\J501671.raw

L:4 M:263 W: Missing Blank Line separator, <120> field identifier
J:8 M:270 C: Current Application Number differs, Replaced Current Application No
L:8 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:24 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1 /
M:332 Repeated in SeqNo=1
L:123 M:254 E: No. of Bases conflict, LENGTH:Input:1440 Counted:1439 SEQ:2 ✓
L:123 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1 ✓
M:254 Repeated in SeqNo=2
L:125 M:252 E: No. of Seq. differs, <211> LENGTH:Input:1548 Found:1547 SEQ:2 ✓
L:136 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:3 differs:8 ✓
L:138 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:8 ✓
M:332 Repeated in SeqNo=3
L:185 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:5 ✓
M:332 Repeated in SeqNo=5
L:292 M:283 W: Missing Blank Line separator, <220> field identifier
L:298 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7 ✓
M:332 Repeated in SeqNo=7
L:349 M:254 E: No. of Bases conflict, LENGTH:Input:12 Counted:22 SEQ:9 ✓
L:349 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2
L:349 M:252 E: No. of Seq. differs, <211> LENGTH:Input:21 Found:22 SEQ:9